## SEOUENCE LISTING

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<110> Tang, Jordan J.N.
     Hong, Lin
      Ghosh, Arun K.
<120> Inhibitors of Memapsin 2 and Use Thereof
<130> OMRF 182
<140> Not Yet Assigned
<141> 2000-06-27
<150> 60/141,363
<151> 1999-06-28
<150> 60/168,060
<151> 1999-11-30
<150> 60/177,836
<151> 2000-01-25
<150> 60/178,368
<151> 2000-01-27
<150> 60/210,292
<151> 2000-06-08
<160> 31
<170> PatentIn Ver. 2.1
<210> 1
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ctggggggcg ccccctggg gctgcggctg ccccgggaga ccgacgaaga gcccgaggag 120
cccggccgga ggggcagctt tgtggagatg gtggacaacc tgaggggcaa gtcggggcag 180
ggctactacg tggagatgac cgtgggcagc ccccgcaga cgctcaacat cctggtggat 240
acaggcagca gtaactttgc agtgggtgct gcccccacc ccttcctgca tcgctactac 300
cagaggcage tgtccagcac ataccgggac ctccggaagg gtgtgtatgt gccctacacc 360
cagggcaagt gggaagggga gctgggcacc gacctggtaa gcatccccca tggccccaac 420
gtcactgtgc gtgccaacat tgctgccatc actgaatcag acaagttctt catcaacggc 480
tccaactggg aaggcatcct ggggctggcc tatgctgaga ttgccaggcc tgacgactcc 540
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ctggagcctt tctttgactc tctggtaaag cagacccacg ttcccaacct cttctccctg 600

cagctttgtg gtgctggctt ccccctcaac cagtctgaag tgctggcctc tgtcggaggg 660 agcatgatca ttggaggtat cgaccactcg ctgtacacag gcagtctctg gtatacaccc 720 atccggcggg agtggtatta tgaggtgatc attgtgcggg tggagatcaa tggacaggat 780 ctgaaaatgg actgcaagga gtacaactat gacaagagca ttgtggacag tggcaccacc 840 aaccttegtt tgeecaagaa agtgtttgaa getgeagtea aateeateaa ggeageetee 900 tecaeggaga agttecetga tggtttetgg etaggagage agetggtgtg etggeaagea 960 ggcaccaccc cttggaacat tttcccagtc atctcactct acctaatggg tgaggttacc 1020 aaccagtect teegeateac cateetteeg cageaatace tgeggeeagt ggaagatgtg 1080 gccacgtccc aagacgactg ttacaagttt gccatctcac agtcatccac gggcactgtt 1140 atgggagctg ttatcatgga gggcttctac gttgtctttg atcgggcccg aaaacgaatt 1200 ggctttgctg tcagcgcttg ccatgtgcac gatgagttca ggacggcagc ggtggaaggc 1260 ccttttgtca ccttggacat ggaagactgt ggctacaaca ttccacagac agatgagtca 1320 acceteatga ceatageeta tgteatgget geeatetgeg ceetetteat getgeeacte 1380 tgcctcatgg tgtgtcagtg gcgctgcctc cgctgcctgc gccagcagca tgatgacttt 1440 gctgatgaca tctccctgct gaagtgagga ggcccatggg cagaagatag agattcccct 1500 ggaccacacc tccgtggttc actttggtca caagtaggag acacagatgg cacctgtggc 1560 cagagcacct caggaccctc cccacccacc aaatgcctct gccttgatgg agaaggaaaa 1620 ggctggcaag gtgggttcca gggactgtac ctgtaggaaa cagaaaagag aagaaagaag 1680 cactetgetg gegggaatac tettggteac etcaaattta agtegggaaa ttetgetget 1740 tgaaacttca gccctgaacc tttgtccacc attcctttaa attctccaac ccaaagtatt 1800 cttcttttct tagtttcaga agtactggca tcacacgcag gttaccttgg cgtgtgtccc 1860 tgtggtaccc tggcagagaa gagaccaagc ttgtttccct gctggccaaa gtcagtagga 1920 gaggatgcac agtttgctat ttgctttaga gacagggact gtataaacaa gcctaacatt 1980 ggtgcaaaga ttgcctcttg aattaaaaaa aaactagatt gactatttat acaaatgggg 2040 gcggctggaa agaggagaag gagagggagt acaaagacag ggaatagtgg gatcaaagct 2100 aggaaaggca gaaacacaac cactcaccag tectagtttt agaceteate tecaagatag 2160 catcccatct cagaagatgg gtgttgtttt caatgttttc ttttctgtgg ttgcagcctg 2220 accaaaagtg agatgggaag ggcttatcta gccaaagagc tcttttttag ctctcttaaa 2280 tgaagtgccc actaagaagt tccacttaac acatgaattt ctgccatatt aatttcattg 2340 tetetatetg aaccaeeett tattetaeat atgataggea geaetgaaat ateetaaeee 2400 cctaagetee aggtgeeetg tgggagagea actggaetat ageagggetg ggetetgtet 2460 teetggteat aggeteacte ttteecceaa atetteetet ggagetttge agceaaggtg 2520 ctaaaaggaa taggtaggag acctetteta tetaateett aaaageataa tgttgaacat 2580 tcattcaaca gctgatgccc tataacccct gcctggattt cttcctatta ggctataaga 2640 agtagcaaga tetttacata atteagagtg gttteattge etteetacee tetetaatgg 2700 eccetecatt tatttgaeta aageaterea eagtggeact ageattatae caagagtatg 2760 agaaatacag tgctttatgg ctctaacatt actgccttca gtatcaaggc tgcctggaga 2820 aaggatggca gcctcagggc ttccttatgt cctccaccac aagagctcct tgatgaaggt 2880 catctttttc ccctatcctg ttcttcccct ccccgctcct aatggtacgt gggtacccag 2940 gctggttctt gggctaggta gtggggacca agttcattac ctccctatca gttctagcat 3000 agtaaactac ggtaccagtg ttagtgggaa gagctgggtt ttcctagtat acccactgca 3060 tectactect acctggteaa eccgetgett ecaggtatgg gacetgetaa gtgtggaatt 3120 acctgataag ggagagggaa atacaaggag ggcctctggt gttcctggcc tcagccagct 3180 gcccmcaagc cataaaccaa taaamcaaga atactgagtc taaaaaaaaaa aaaaaaaaa 3240 aaaaaaaaa aa 3252 <211> 488

<212> PRT

<213> Homo sapiens

<220>

<223> Purified Memapsin 2

<220>

<223> Amino Acids 28-48 are remnant putative propeptide residues

<220>

<223> Amino Acids 58-61, 78, 80, 82-83, 116, 118-121,
 156, 166, 174, 246, 274, 276, 278-281, 283, and
 376-377 are residues in contact with the OM99-2
 inhibitor

<220>

<223> Amino acids 54-57, 61-68, 73-80, 86-89, 109-111, 113-118, 123-134, 143-154, 165-168, 198-202, and 220-224 are N-lobe Beta Strands

<220>

<223> Amino Acids 184-191 and 210-217 are N-lobe Helices

<220>

<220>

<223> Amino Acids 286-299, 307-310, 350-353, 384-387, and 427-431 are C-lobe Helices

<400> 2

Ala Gly Val Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro

1 5 10 15

Leu Arg Ser Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg
20 25 30

Glu Thr Asp Glu Glu Pro Glu Glu Pro Gly Arg Arg Gly Ser Phe Val
35 40 45

Glu Met Val Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val
50 55 60

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						•										
	Glu 65	Met	Thr	Val	Gly	Ser 70	Pro	Pro	Gln	Thr	Leu 75	Asn	Ile	Leu	Val	Asp 80
	Thr	Gly	Ser	Ser	Asn 85	Phe	Ala	Val	Gly	Ala 90	Ala	Pro	His	Pro	Phe 95	Leu
	His	Arg	Tyr	Tyr 100	Gln	Arg	Gln	Leu	Ser 105	Ser	Thr	Tyr	Arg	Asp 110	Leu	Arg
	Lys	Gly	Val 115	Tyr	Val	Pro	Tyr	Thr 120	Gln	Gly	Lys	Trp	Glu 125	Gly	Glu	Leu
	Gly	Thr 130	Asp	Leu	Val	Ser	Ile 135	Pro	His	Gly	Pro	Asn 140	Val	Thr	Val	Arg
	Ala 145	Asn	Ile	Ala	Ala	Ile 150	Thr	Glu	Ser	Asp	Lys 155	Phe	Phe	Ile	Asn	Gly 160
	Ser	Asn	Trp	Glu	Gly 165	Ile	Leu	Gly	Leu	Ala 170	Tyr	Ala	Glu	Ile	Ala 175	Arg
	Pro	Asp	Asp	Ser 180	Leu	Glu	Pro	Phe	Phe 185	Asp	Ser	Leu	Val	Lys 190	Gln	Thr
	His	Val	Pro 195	Asn	Leu	Phe	Ser	Leu 200	Gln	Leu	Cys	Gly	Ala 205	Gly	Phe	Pro
	Leu	Asn 210	Gln	Ser	Glu	Val	Leu 215	Ala	Ser	Val	Gly	Gly 220	Ser	Met	Ile	Ile
	Gly 225	Gly	Ile	Asp	His	Ser 230	Leu	Tyr	Thr	Gly	Ser 235	Leu	Trp	Tyr	Thr	Pro 240
	Ile	Arg	Arg	Glu	Trp 245	туг	Tyr	Glu	Val	Ile 250	Ile	Val	Arg	Val	Glu 255	Ile
	Asn	Gly	Gln	Asp 260	Leu	Lys	Met	Asp	Cys 265	Lys	Glu	Tyr	Asn	Tyr 270	Asp	Lys
	Ser	Ile	Val 275	Asp	Ser	Gly	Thr	Thr 280	Asn	Leu	Arg	Leu	Pro 285	Lys	Lys	Val
	Phe	Glu 290	Ala	Ala	Val	Lys	Ser 295	Ile	Lys	Ala	Ala	Ser 300	Ser	Thr	Glu	Lys
	Phe 305	Pro	Asp	Gly	Phe	Trp 310	Leu	Gly	Glu	Gln	Leu 315	Val	Cys	Trp	Gln	Ala 320

Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser Leu Tyr Leu Met 325 330 Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu Pro Gln Gln 340 345 Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp Asp Cys Tyr 360 Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val Met Gly Ala Val 375 Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg Lys Arg Ile 390 395 Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu Phe Arg Thr Ala 405 410 Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu Asp Cys Gly Tyr 420 425 430 Asn Ile Pro Gln Thr Asp Glu Ser Thr Leu Met Thr Ile Ala Tyr Val 435 440 Met Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val 450 455 Cys Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe 465 470 475 480

Ala Asp Asp Ile Ser Leu Leu Lys . 485

<210>. 3

<211> 503

<212> PRT

<213> Homo sapiens

<220>

<223> Pro-memapsin 2

<220>

<223> Amino Acids 1-15 are vector-derived residues

<220>

<223> Amino Acids 16-64 are a putative pro peptide

<220>

<223> Amino Acids 1-13 are the T7 promoter

<220>

<223> Amino Acids 16-456 are Pro-memapsin 2-T1

<220>

<223> Amino Acids 16-421 are Promemapsin 2-T2

<400> 3

Met Ala Ser Met Thr Gly Gly Gln Gln Met Gly Arg Gly Ser Met Ala

1 5 10 15

Gly Val Leu Pro Ala His Gly Thr Gln His Gly Ile Arg Leu Pro Leu 20 25 30

Arg Ser Gly Leu Gly Gly Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu
35 40 45

Thr Asp Glu Glu Pro Glu Pro Gly Arg Arg Gly Ser Phe Val Glu
50 55 60

Met Val Asp Asn Leu Arg Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu 65 70 75 80

Met Thr Val Gly Ser Pro Pro Gln Thr Leu Asn Ile Leu Val Asp Thr
85 90 95

Gly Ser Ser Asn Phe Ala Val Gly Ala Ala Pro His Pro Phe Leu His
100 105 110

Arg Tyr Tyr Gln Arg Gln Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys 115 120 125

Gly Val Tyr Val Pro Tyr Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly
130 135 140

Thr Asp Leu Val Ser Ile Pro His Gly Pro Asn Val Thr Val Arg Ala 145 150 155 160

Asn Ile Ala Ala Ile Thr Glu Ser Asp Lys Phe Phe Ile Asn Gly Ser 165 170 175

Asn Trp Glu Gly Ile Leu Gly Leu Ala Tyr Ala Glu Ile Ala Arg Pro 180 185 190

Asp Asp Ser Leu Glu Pro Phe Phe Asp Ser Leu Val Lys Gln Thr His 195 200 205

val	210	ASII	Leu	PHE	ser	215	GIN	ren	Сув	GIY	220	GIĀ	Pne	PIO	Leu
Asn 225	Gln	Ser	Glu	Val	Leu 230	Ala	Ser	Val	Gly	Gly 235	Ser	Met	Ile	Ile	Gly 240
Gly	Ile	Asp	His	Ser 245	Leu	Tyr	Thr	Gly	Ser 250	Leu	Trp	Tyr	Thr	Pro 255	Ile
Arg	Arg	Glu	Trp 260	Tyr	Tyr	Glu	Val	Ile 265	Ile	Val	Arg	Val	Glu 270	Ile	Asn
Gly	Gln	Asp 275	Leu	Lys	Met	Asp	Cys 280	Lys	Glu	Tyr	Asn	Tyr 285	Asp	Lys	Ser
Ile	Val 290	Asp	Ser	Gly	Thr	Thr 295	Asn	Leu	Arg	Leu	Pro 300	Lys	Lys	Val	Phe
305					310	Ile	_			315					320
٠				325		Gly			330					335	
			340			Phe		345				-	350		_
		355				Phe	360	•				365			
	370				_	Val 375					380	_	_	_	_
385					390	Ser				395					400
•				405		Val			410					415	
			420		_	His		425	_				430		
		435				Thr	440					445			
Ile	Pro 450	Gln	Thr	Asp	Glu	Ser 455	Thr	Leu	Met	Thr	11e 460	Ala	Tyr	Val	Met

```
Ala Ala Ile Cys Ala Leu Phe Met Leu Pro Leu Cys Leu Met Val Cys
465
                    470
                                         475
                                                              480
Gln Trp Arg Cys Leu Arg Cys Leu Arg Gln Gln His Asp Asp Phe Ala
                485
                                     490
Asp Asp Ile Ser Leu Leu Lys
          500
<210> 4
<211> 10
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 4
Ser Glu Val Lys Met Asp Ala Glu Phe Arg
<210> 5
<211> 10
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 5
Ser Glu Val Asn Leu Asp Ala Glu Phe Arg
  1
<210> 6
<211> 8
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 6

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Ser Val Asn Met Ala Glu Gly Asp
<210> 7
<211> 12
<212> PRT
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      Peptide
<400> 7
Lys Gly Gly Val Val Ile Ala Thr Val Ile Val Lys
                  5
                                      10
<210> 8
<211> 4
<212> PRT
<213> Homo sapiens
<400> 8
Asp Thr Ser Gly
 1
<210> 9
<211> 8
<212> PRT
<213> Homo sapiens
<400> 9
Leu Val Asn Met Ala Glu Gly Asp
 1
<210> 10
<211> 28
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 10
```

ggtaagcatc ccccatggcc ccaacgtc

```
<210> 11
<211> 28
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 11
gacgttgggg ccatggggga tgcttacc
                                                                    28
<210> 12
<211> 34
<212> DNA
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<220>
<223> Description of Artificial Sequence: Primer
<400> 12
acgttgtctt tgatcgggcc cgaaaacgaa ttgg
                                                                    34
<210> 13
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 13
ccaattcgtt ttcgggcccg atcaaagaca acg
                                                                   33
<210> 14
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 14
ccatcctaat acgactcact atagggc
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27

```
<210> 15
<211> 23
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 15
actcactata gggctcgagc ggc
                                                                    23
<210> 16
<211> 26
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 16
cttttgagca agttcagcct ggttaa
                                                                    26
<210> 17
<211> 31
<212> DNA
<213> Artificial Sequence
<223> Description of Artificial Sequence: Primer
<400> 17
gaggtggctt atgagtattt cttccagggt a
                                                                    31
<210> 18
<211> 22
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 18
tggcgacgac tcctggagcc cg
                                                                    22
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```
<210> 19
<211> 24
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 19
tgacaccaga ccaactggta atgg
                                                                    24
<210> 20
<211> 27
<212> DNA
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Primer
<400> 20
                                                                    27
catatggcgg gagtgctgcc tgcccac
<210> 21
<211> 38
<212> DNA
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Primer
<400> 21
                                                                    38
ggatcctcac ttcagcaggg agatgtcatc agcaaagt
<210> 22
<211> 8
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<220>
<223> Description of Artificial Sequence: Oxidized
      Insulin B-chain
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<220>

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<223> Xaa at site 3 represents cysteic acid
<400> 22
His Leu Xaa Gly Ser His Leu Val
<210> 23
<211> 8
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Oxidized
      Insulin B-chain
<220>
<223> Xaa at site 1 represents cysteic acid
<400> 23
Xaa Gly Glu Arg Gly Phe Phe Tyr
<210> 24
<211> 5
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 24
Val Gly Ser Gly Val
  1
<210> 25
<211> 7
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
      Peptide
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<400> 25

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Val Gly Ser Gly Val Leu Leu
  1
<210> 26
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Synthetic
      Peptide
<400> 26
Gly Val Leu Leu Ser Arg Lys
<210> 27
<211> 7
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Inhibitors
<400> 27
Val Asn Leu Ala Ala Glu Phe
  1
                  5
<210> 28
<211> 8
<212> PRT
<213> Artificial Sequence
<223> Description of Artificial Sequence: Inhibitors
<400> 28
Glu Val Asn Leu Ala Ala Glu Phe
  1
<210> 29
<211> 4
<212> PRT
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<213> Artificial Sequence

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<220>
 <223> Description of Artificial Sequence: Synthetic
       Peptide
 <400> 29
 Asn Leu Ala Ala
   1
 <210> 30
 <211> 10
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       Peptide
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Val Gly Ser Gly Val Leu Leu Ser Arg Lys
   1
                   5
 <210> 31
 <211> 326
 <212> PRT
 <213> Homo sapiens
 <220>
 <223> Amino acids 2-5, 6-9, 13-20, 25-32, 65-67, 69-74,
       79-87, 89-91, 99-106, 119-122, 150-154, 164-167,
       180-183, 191-194, 196-199, 201-204, 210-214,
       221-223, 258-262, 265-269, and 275-278 are Beta
       Strands
 <220>
 <223> Amino acids 281-284, 286-288, 298-301, 310-315,
       and 319-324 are Beta strands
 <220>
 <223> Amino acids 48-51, 111-114, 136-142, 225-234,
       249-254, 271-274, and 303-306 are Helices
 <220>
 <223> Amino acids 12-13, 30, 32, 34-35, 73-77, 111, 117,
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13 4 E

120, 189, 213, 215, 217-220, 287, 289, 291, 298, and 300 are residues in contact with pepstatin.

<220>

<223> Pepsin

<400> 31

Val Asp Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly

1 5 10 15

Thr Ile Gly Ile Gly Thr Pro Ala Gln Asp Phe Thr Val Val Phe Asp 20 25 30

Thr Gly Ser Ser Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu 35 40 45

Ala Cys Thr Asn His Asn Arg Phe Asn Pro Glu Asp Ser Ser Thr Tyr 50 55 60

Gln Ser Thr Ser Glu Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met
65 70 75 80

Thr Gly Ile Leu Gly Tyr Asp Thr Val Gln Val Gly Gly Ile Ser Asp
85
90
95

Thr Asn Gln Ile Phe Gly Leu Ser Glu Thr Glu Pro Gly Ser Phe Leu
100 105 110

Tyr Tyr Ala Pro Phe Asp Gly Ile Leu Gly Leu Ala Tyr Pro Ser Ile 115 120 125

Ser Ser Ser Gly Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly 130 135 140

Leu Val Ser Gln Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln 145 150 155 160

Ser Gly Ser Val Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr 165 170 175

Gly Ser Leu Asn Trp Val Pro Val Thr Val Glu Gly Tyr Trp Gln Ile 180 185 190

Thr Val Asp Ser Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu 195 200 205

Gly Cys Gln Ala Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Gly Pro 210 215 220

Thr Ser Pro Ile Ala Asn Ile Gln Ser Asp Ile Gly Ala Ser Glu Asn

Ser Asp Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro 245 250 255

Asp Ile Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser 260 265 270

Ala Tyr Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly
275 280 285

Met Asn Leu Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val 290 295 300

Phe Ile Arg Gln Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn Gln Val 305 310 315

Gly Leu Ala Pro Val Ala 325